

#8



PCT09

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/936,205

DATE: 02/08/2002  
TIME: 11:26:41

Input Set : A:\37945-0024.txt  
Output Set: N:\CRF3\02082002\I936205.raw

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4 <110> APPLICANT: SMITH, Richard Anthony Godwin
5      PRATT, Julian Roy
6      SACKS, Steven Howard
8 <120> TITLE OF INVENTION: ORGAN TRANSPLANT SOLUTIONS CONTAINING CONJUGATES OF
9      SOLUBLE PEPTIDIC COMPOUNDS WITH MEMBRANE-BINDING
11 <130> FILE REFERENCE: 37945-0024
13 <140> CURRENT APPLICATION NUMBER: US 09/936,205
C--> 14 <141> CURRENT FILING DATE: 2001-12-07
16 <150> PRIOR APPLICATION NUMBER: PCT/GB00/00834
17 <151> PRIOR FILING DATE: 2000-03-08
19 <150> PRIOR APPLICATION NUMBER: GB 9905503.0
20 <151> PRIOR FILING DATE: 1999-03-10
22 <160> NUMBER OF SEQ ID NOS: 11
24 <170> SOFTWARE: PatentIn Ver. 2.1
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 215
28 <212> TYPE: PRT
29 <213> ORGANISM: Artificial Sequence
31 <220> FEATURE:
32 <223> OTHER INFORMATION: Linear, 2 polypeptide chains disulphide linked
34 <220> FEATURE:
35 <221> NAME/KEY: DISULFID
36 <222> LOCATION: (198)..(199)
38 <220> FEATURE:
39 <223> OTHER INFORMATION: 2nd polypeptide chain (199-215) runs C to N
40      terminus
42 <220> FEATURE:
43 <223> OTHER INFORMATION: An N-myristoyl group is at the N-terminus of the
44      second polypeptide chain
46 <220> FEATURE:
47 <223> OTHER INFORMATION: A CONH2 group is at the C terminus of the second
48      polypeptide chain
50 <220> FEATURE:
51 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
52      peptide reagent
54 <400> SEQUENCE: 1
55 Met Gln Cys Asn Ala Pro Glu Trp Leu Pro Phe Ala Arg Pro Thr Asn
56      1           5           10          15
58 Leu Thr Asp Glu Phe Glu Pro Ile Gly Thr Tyr Leu Asn Tyr Glu
59      20          25          30
61 Cys Arg Pro Gly Tyr Ser Gly Arg Pro Phe Ser Ile Ile Cys Leu Lys
62      35          40          45
64 Asn Ser Val Trp Thr Gly Ala Lys Asp Arg Cys Arg Arg Lys Ser Cys

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65 50 55 60  
67 Arg Asn Pro Pro Asp Pro Val Asn Gly Met Val His Val Ile Lys Gly  
68 65 70 75 80  
70 Ile Gln Phe Gly Ser Gln Ile Lys Tyr Ser Cys Thr Lys Gly Tyr Arg  
71 85 90 95  
73 Leu Ile Gly Ser Ser Ser Ala Thr Cys Ile Ile Ser Gly Asp Thr Val  
74 100 105 110  
76 Ile Trp Asp Asn Glu Thr Pro Ile Cys Asp Arg Ile Pro Cys Gly Leu  
77 115 120 125  
79 Pro Pro Thr Ile Thr Asn Gly Asp Phe Ile Ser Thr Asn Arg Glu Asn  
80 130 135 140  
82 Phe His Tyr Gly Ser Val Val Thr Tyr Arg Cys Asn Pro Gly Ser Gly  
83 145 150 155 160  
85 Gly Arg Lys Val Phe Glu Leu Val Gly Glu Pro Ser Ile Tyr Cys Thr  
86 165 170 175  
88 Ser Asn Asp Asp Gln Val Gly Ile Trp Ser Gly Pro Ala Pro Gln Cys  
89 180 185 190  
91 Ile Ile Pro Asn Lys Cys Cys Asp Gly Pro Lys Lys Lys Lys Lys Lys  
92 195 200 205  
94 Ser Pro Ser Lys Ser Ser Gly  
95 210 215  
98 <210> SEQ ID NO: 2  
99 <211> LENGTH: 218  
100 <212> TYPE: PRT  
101 <213> ORGANISM: Artificial Sequence  
103 <220> FEATURE:  
104 <223> OTHER INFORMATION: 2 polypeptide chains disulphide linked  
106 <220> FEATURE:  
107 <221> NAME/KEY: DISULFID  
108 <222> LOCATION: (198)..(199)  
110 <220> FEATURE:  
111 <223> OTHER INFORMATION: The second polypeptide chain (199-218) runs C to N  
112 terminus  
114 <220> FEATURE:  
115 <223> OTHER INFORMATION: An N-Myristoyl group is at the N terminus of the  
116 second polypeptide chain  
118 <220> FEATURE:  
119 <223> OTHER INFORMATION: A CONH<sub>2</sub> group is at the C terminus of the second  
120 polypeptide chain  
122 <220> FEATURE:  
123 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic  
124 peptide reagent  
126 <400> SEQUENCE: 2  
127 Met Gln Cys Asn Ala Pro Glu Trp Leu Pro Phe Ala Arg Pro Thr Asn  
128 1 5 10 15  
130 Leu Thr Asp Glu Phe Glu Phe Pro Ile Gly Thr Tyr Leu Asn Tyr Glu  
131 20 25 30  
133 Cys Arg Pro Gly Tyr Ser Gly Arg Pro Phe Ser Ile Ile Cys Leu Lys  
134 35 40 45

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136 Asn Ser Val Trp Thr Gly Ala Lys Asp Arg Cys Arg Arg Lys Ser Cys  
137 50 55 60  
139 Arg Asn Pro Pro Asp Pro Val Asn Gly Met Val His Val Ile Lys Gly  
140 65 70 75 80  
142 Ile Gln Phe Gly Ser Gln Ile Lys Tyr Ser Cys Thr Lys Gly Tyr Arg  
143 85 90 95  
145 Leu Ile Gly Ser Ser Ser Ala Thr Cys Ile Ile Ser Gly Asp Thr Val  
146 100 105 110  
148 Ile Trp Asp Asn Glu Thr Pro Ile Cys Asp Arg Ile Pro Cys Gly Leu  
149 115 120 125  
151 Pro Pro Thr Ile Thr Asn Gly Asp Phe Ile Ser Thr Asn Arg Glu Asn  
152 130 135 140  
154 Phe His Tyr Gly Ser Val Val Thr Tyr Arg Cys Asn Pro Gly Ser Gly  
155 145 150 155 160  
157 Gly Arg Lys Val Phe Glu Leu Val Gly Glu Pro Ser Ile Tyr Cys Thr  
158 165 170 175  
160 Ser Asn Asp Asp Gln Val Gly Ile Trp Ser Gly Pro Ala Pro Gln Cys  
161 180 185 190  
163 Ile Ile Pro Asn Lys Cys Cys Ala Asp Leu Arg Ser Ser Leu Gly Pro  
164 195 200 205  
166 Lys Lys Lys Lys Lys Ser Pro Ser Gly  
167 210 215  
170 <210> SEQ ID NO: 3  
171 <211> LENGTH: 20  
172 <212> TYPE: PRT  
173 <213> ORGANISM: Artificial Sequence  
175 <220> FEATURE:  
176 <223> OTHER INFORMATION: An N-myristoyl group is at the N terminus of the  
177 polypeptide chain  
179 <220> FEATURE:  
180 <223> OTHER INFORMATION: A CONH<sub>2</sub> group is at the C-terminus of the  
181 polypeptide chain  
183 <220> FEATURE:  
184 <223> OTHER INFORMATION: An S-2-Thiopyridyl group is attached to the  
185 C-terminal cysteine  
187 <220> FEATURE:  
188 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic  
189 peptide reagent  
191 <400> SEQUENCE: 3  
192 Gly Ser Pro Ser Lys Lys Lys Lys Lys Pro Gly Leu Ser Ser Arg  
193 1 5 10 15  
195 Leu Asp Ala Cys  
196 20  
199 <210> SEQ ID NO: 4  
200 <211> LENGTH: 20  
201 <212> TYPE: PRT  
202 <213> ORGANISM: Artificial Sequence  
204 <220> FEATURE:  
205 <223> OTHER INFORMATION: A CONH<sub>2</sub> group is at the C terminus

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207 <220> FEATURE:  
208 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic  
209 peptide  
211 <400> SEQUENCE: 4  
212 Gly Ser Pro Ser Lys Lys Lys Lys Lys Pro Gly Leu Ser Ser Arg  
213 1 5 10 15  
215 Leu Asp Ala Cys  
216 20  
219 <210> SEQ ID NO: 5  
220 <211> LENGTH: 9  
221 <212> TYPE: PRT  
222 <213> ORGANISM: Artificial Sequence  
224 <220> FEATURE:  
225 <223> OTHER INFORMATION: Description of Artificial Sequence: A peptidic  
226 membrane binding element of SEQ ID NO: 4  
228 <400> SEQUENCE: 5  
229 Pro Ser Lys Lys Lys Lys Lys Pro  
230 1 5  
233 <210> SEQ ID NO: 6  
234 <211> LENGTH: 7  
235 <212> TYPE: PRT  
236 <213> ORGANISM: Artificial Sequence  
238 <220> FEATURE:  
239 <223> OTHER INFORMATION: Description of Artificial Sequence: A peptidic  
240 membrane binding element of SEQ ID NO: 4  
242 <400> SEQUENCE: 6  
243 Leu Ser Ser Arg Leu Asp Ala  
244 1 5  
247 <210> SEQ ID NO: 7  
248 <211> LENGTH: 16  
249 <212> TYPE: PRT  
250 <213> ORGANISM: Artificial Sequence  
252 <220> FEATURE:  
253 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of  
254 electrostatic switch sequence  
256 <400> SEQUENCE: 7  
257 Asp Gly Pro Lys Lys Lys Lys Lys Ser Pro Ser Lys Ser Ser Gly  
258 1 5 10 15  
261 <210> SEQ ID NO: 8  
262 <211> LENGTH: 16  
263 <212> TYPE: PRT  
264 <213> ORGANISM: Artificial Sequence  
266 <220> FEATURE:  
267 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of  
268 electrostatic switch sequence  
270 <400> SEQUENCE: 8  
271 Gly Ser Ser Lys Ser Pro Ser Lys Lys Lys Lys Lys Pro Gly Asp  
272 1 5 10 15  
275 <210> SEQ ID NO: 9

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276 <211> LENGTH: 20  
277 <212> TYPE: PRT  
278 <213> ORGANISM: Artificial Sequence  
280 <220> FEATURE:  
281 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of  
282 electrostatic switch sequence  
284 <400> SEQUENCE: 9  
285 Ser Pro Ser Asn Glu Thr Pro Lys Lys Lys Lys Arg Phe Ser Phe  
286 1 5 10 15  
288 Lys Lys Ser Gly  
289 20  
292 <210> SEQ ID NO: 10  
293 <211> LENGTH: 16  
294 <212> TYPE: PRT  
295 <213> ORGANISM: Artificial Sequence  
297 <220> FEATURE:  
298 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of  
299 electrostatic switch sequence  
301 <400> SEQUENCE: 10  
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303 1 5 10 15  
306 <210> SEQ ID NO: 11  
307 <211> LENGTH: 14  
308 <212> TYPE: PRT  
309 <213> ORGANISM: Artificial Sequence  
311 <220> FEATURE:  
312 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of  
313 electrostatic switch sequence  
315 <400> SEQUENCE: 11  
316 Ser Lys Asp Gly Lys Lys Lys Lys Lys Ser Lys Thr Lys  
317 1 5 10

**VERIFICATION SUMMARY**  
PATENT APPLICATION: US/09/936,205

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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date